Listing of Claims:

- 1. (previously presented) An ApoA-I agonist compound comprising:
- (i) a 15 to 26- residue peptide or peptide analogue according to formula (I) which forms an amphipathic α -helix in the presence of lipids and exhibits at least about 38% LCAT activation activity as compared with human ApoA-I wherein one or two helical turns are deleted from formula (I), wherein a helical turn consists of 3 to 4 consecutive residues selected from residues X_1 to X_{23} of formula (I):

 $Z_1-X_1-X_2-X_3-X_4-X_5-X_6-X_7-X_8-X_9-X_{10}-X_{11}-X_{12}-X_{13}-X_{14}-X_{15}-X_{16}-X_{17}-X_{18}-X_{19}-X_{20}-X_{21}-X_{22}-X_{23}-Z_2$ or a pharmaceutically acceptable salt thereof, wherein:

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X<sub>1</sub> is Pro (P), Ala (A), Gly (G), Gln (Q), Asn (N), Asp (D) or D-Pro (p);
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 X_2 is an aliphatic residue;

 X_3 is a Leu (L) or Phe (F);

 X_4 is Glu (E)

 X_5 is an aliphatic residue;

 X_6 is Leu (L) or Phe (F);

 X_7 is Glu (E) or Leu (L);

 X_8 is Asn (N) or Gln (Q);

 X_9 is Leu (L);

 X_{10} is Leu (L), Trp (W) or Gly (G);

 X_{11} is an acidic residue;

 X_{12} is Arg (R);

 X_{13} is Leu (L) or Gly (G);

 X_{14} is Leu (L), Phe (F) or Gly (G);

 X_{15} is Asp (D);

 X_{16} is Ala (A);

 X_{17} is Leu (L);

 X_{18} is Asn (N) or Gln (Q);

 X_{19} is a basic residue;

 X_{20} is a basic residue;

 X_{21} is Leu (L);

 X_{22} is a basic residue;

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 X_{23} is absent or a basic residue;

- Z_1 is H_2N_- ;
- Z_2 is -C (O) NRR or -C (O) OR;

each R is independently -H, (C_1-C_6) alkyl, (C_1-C_6) alkenyl, (C_1-C_6) alkynyl, (C_5-C_{20}) aryl, (C_6-C_{26}) alkaryl, 5-20 membered heteroaryl or 6-26 membered alkheteroaryl or a 1 to 7-residue peptide or peptide analogue in which one more bonds between residues 1-7 are independently a substituted amide, an isostere of an amide or an amide mimetic; and

each "-" between residues X_1 to X_{23} and between residues of the peptide to Z_2 independently designates an amide linkage, a substituted amide linkage, an isostere of an amide or an amide mimetic; or

an N- terminally blocked form, a C-terminally blocked form, or an N- and C-terminally blocked form of formula (I).

2-55. (canceled)

- 56. (previously presented) The 15 to 26-residue peptide or deleted peptide analogue of Claim 1, in which one helical turn is deleted.
- 57. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which three, four, six, seven or eight residues X_1 , X_2 , X_3 , X_4 , X_5 , X_6 , X_7 , X_8 , X_9 , X_{10} , X_{11} , X_{12} , X_{13} , X_{14} , X_{15} , X_{16} , X_{17} , X_{18} , X_{19} , X_{20} , X_{21} and X_{22} are deleted.
- 58. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 3 consecutive residues are deleted.
- 59. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 4 consecutive residues are deleted.
- 60. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which two non-contiguous sets of 3 consecutive residues are deleted.
- 61. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which two non-contiguous sets of 4 consecutive residues are deleted.

- 62. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which one set of 3 consecutive residues and one set of 4 consecutive residues are deleted.
- 63. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 57, in which 6, 7 or 8 consecutive residues are deleted.

64-66. (canceled)

67. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1 in which:

the "-" between residues designates -C (O) NH-; Z_1 is H_2N -; and Z_2 is -C (O) OH or a salt thereof.

- 68. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobic moment, $\langle \mu_{\rm H} \rangle$, is 0.45 to 0.65.
- 69. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 68, in which the mean hydrophobic moment, $\langle \mu_{\rm H} \rangle$, is 0.50 to 0.60.
- 70. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity, $\langle H_o \rangle$, is -0.050 to -0.070.
- 71. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity, $\langle H_o \rangle$, is -0.030 to -0.055.
- 72. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the mean hydrophobicity of the hydrophobic face, $\langle H_o^{pho} \rangle$, is 0.90 to 1.20.
- 73. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 72, in which the mean hydrophobicity of the hydrophobic face, $\langle H_o^{pho} \rangle$, is 0.94 to 1.10.
- 74. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 1, in which the pho angle is 160° to 220°.

75. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 74, in which the pho angle is 180° to 200°.

76-78. (canceled)

79. (previously presented) A pharmaceutical composition comprising an ApoA-I agonist compound and a pharmaceutically acceptable carrier, excipient or diluent, wherein the ApoA-I agonist compound is a 15 to 26-residue peptide or peptide analogue according to Claim 1 or 57.

80-81. (canceled)

- 82. (previously presented) The pharmaceutical composition of Claim 79 which is a lyophilized powder.
- 83. (previously presented) The pharmaceutical composition of Claim 79 which is a solution.
- 84. (previously presented) The N-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.
- 85. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 84 in which the N-terminally blocking group is selected from the group consisting of acetyl, formyl and dansyl.
- 86. (previously presented) The C-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.
- 87. (previously presented) The 15 to 26-residue peptide or peptide analogue of Claim 86 in which the C-terminally blocking group is methyl.
- 88. (previously presented) The N-terminally and C-terminally blocked form of the 15 to 26-residue peptide or peptide analogue of Claim 1.